

17

62

107

152

197

242

260

310

360

410

431

Fig. 1

00450-444E60

Met	Glu	Thr	Arg	Ala	Phe	Trp	Ile	Thr	Leu	Leu	Leu	Val	Leu	Val
				5					10				15	
Ala	Gly	Ser	Ser	Cys	Lys	Ala	Gln	Glu	Phe	Val	Gly	Leu	Ser	Pro
				20					25				30	
Ser	Gln	Cys	Met	Ala	Pro	Thr	Asn	Val	Arg	Val	Asp	Cys	Asn	Tyr
				35					40				45	
Pro	Thr	Val	Thr	Ser	Glu	Gln	Cys	Asn	Asn	Arg	Gly	Cys	Cys	Phe
				50					55				60	
Asp	Ser	Ser	Ile	Pro	Asn	Val	Pro	Trp	Cys	Phe	Lys	Pro	Leu	Gln
				65					70				75	
Glu	Thr	Glu	Cys	Thr	Phe									
				80										

Fig 2

rITF METRAFWITLLLVLVAGSSCKAQEFVGLSPSQCMAPTNVRVDCNYPTVTSEQCNNRGCC
 ps2 -----EAQ-----TETCTVAPRERQNCGFPGVTPSQCANKGCC
 PSP -----EKPAACRCSRQDPKN-RVNCGFPGITSDQCFTSGCC
 rITF Loop 3 = 10 FDSSIPNVPWCFK-----PLQ-----ETECT-----F
 ps2 FDDTVRGVPWCFY-----PNTIDVPPEEECE-----F
 PSP FDSQVPGVPWCFK-----PLP-----AQESEECVMEV

Fig. 3

Y2610593 11/19/69 10:00 AM
RECEIVED 11/19/69 10:00 AM
U.S. AIR FORCE
HONOLULU, HAWAII

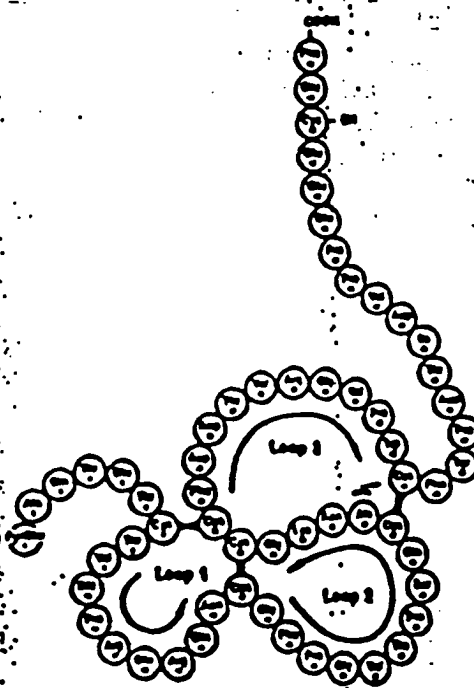


Fig. 4A

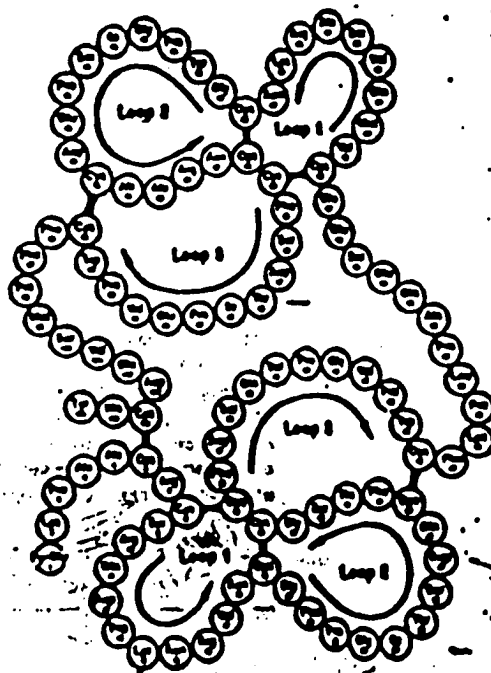


Fig. 4B

66-150-444-1-60

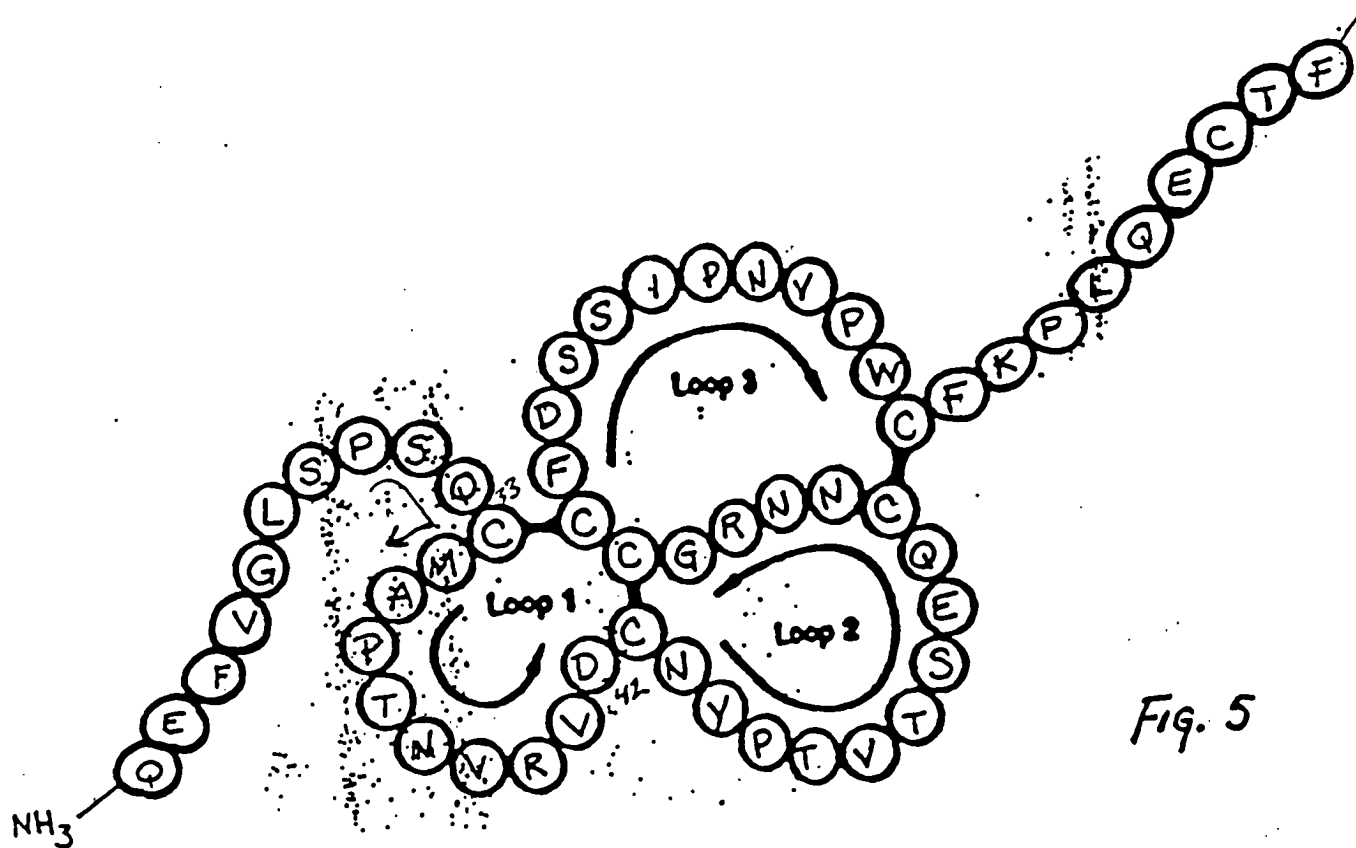


Fig. 5

g atg ctg ggg ctg gtc ctg gcc ttg ctg tcc tcc agc tct gct gag gag	49
Met Leu Gly Leu Val Leu Ala Leu Leu Ser Ser Ser Ser Ala Glu Glu	
1 5 10 15	
tac gtg ggc ctg tct gca aac cag tgt gcc gtg ccg gcc aag gac agg	97
Tyr Val Gly Leu Ser Ala Asn Gln Cys Ala Val Pro Ala Lys Asp Arg	
20 25 30	
gtg gac tgc ggc tac ccc cat gtc acc ccc aag gag tgc aac aac cgg	145
Val Asp Cys Gly Tyr Pro His Val Thr Pro Lys Glu Cys Asn Asn Arg	
35 40 45	
ggc tgc tgc ttt gac tcc agg atc cct gga gtg cct tgg tgt ttc aag	193
Gly Cys Cys Phe Asp Ser Arg Ile Pro Gly Val Pro Trp C P K	
50 55 60	
ccc ctg cag gaa gca gaa tgc acc ttc tgaggcacct ccagctgccc	243
P L Q E A E C T F	
65 70	
ctgggatgca ggctgagcac ccttgcccgg ctgtgattgc tgccaggcac tgttcatctc	303
agtttttctg tccctttgct cccggcaagc tttctgctga aagttcatat ctggagcctg	363
atgtcttaac gaataaaggt cccatgctcc acccgaaaaa	403

FIG. 6

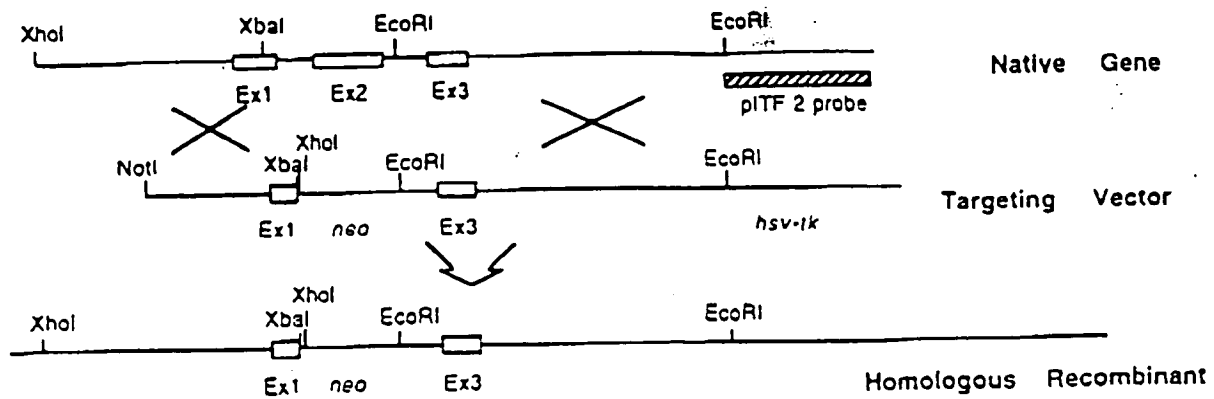


Fig. 7

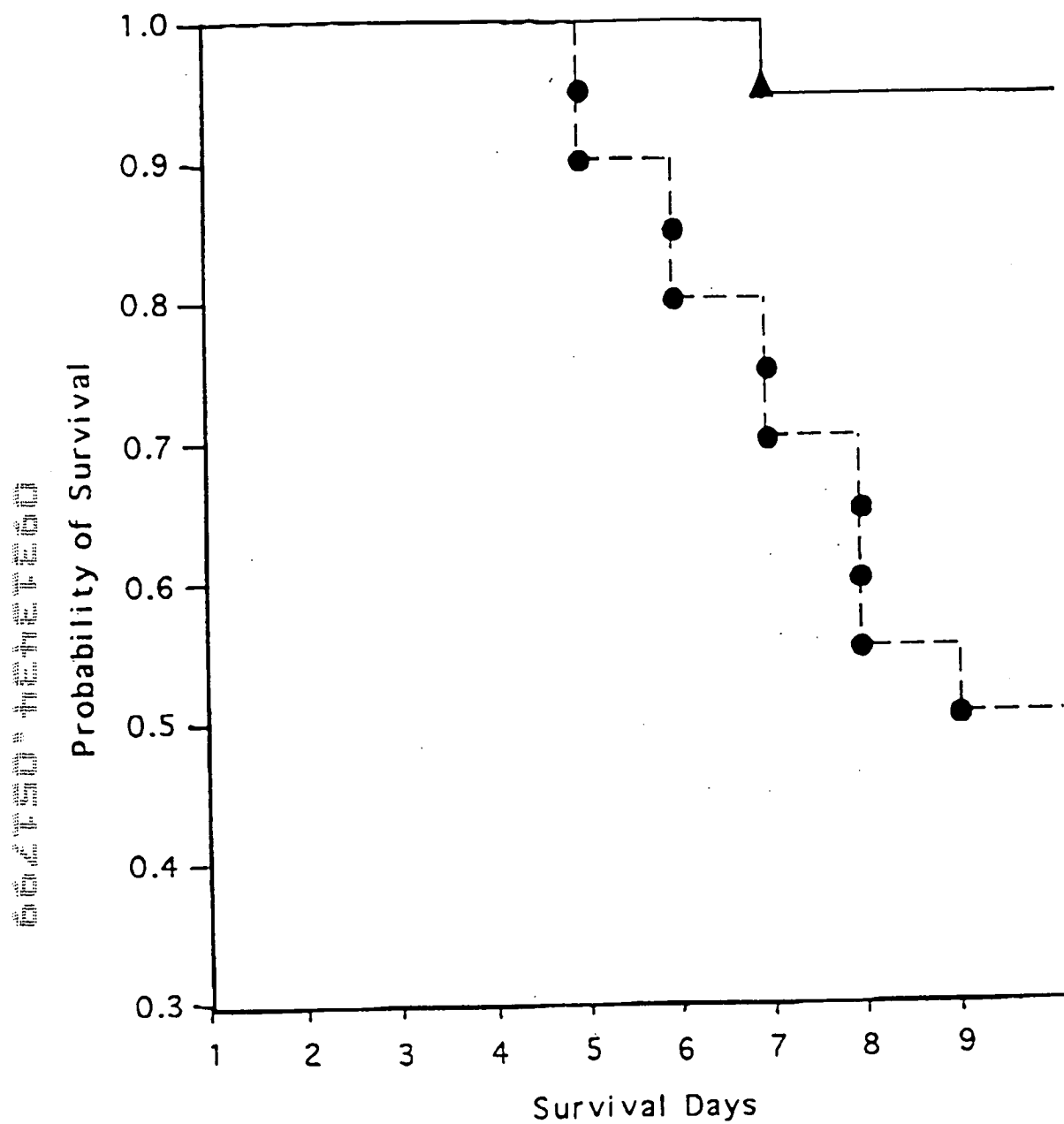


Fig. 8

FIG. 9

atccctgact	cggggtcgcc	tttgagcag	agaggaggca	atg gcc acc atg gag	55
				Met Ala Thr Met Glu	
				1 5	
aac aag gtg atc tgc gcc ctg gtc ctg gtg tcc atg ctg gcc ctc ggc	103				
Asn Lys Val Ile Cys Ala Leu Val Leu Val Ser Met Leu Ala Leu Gly					
	10 15 20				
acc ctg gcc gag gcc cag aca gag acg tgt aca gtg gcc ccc cgt gaa	151				
Thr Leu Ala Glu Ala Gln Thr Glu Thr Cys Thr Val Ala Pro Arg Glu					
	25 30 35				
aga cag aat tgt ggt ttt cct ggt gtc acg ccc tcc cag tgt gca aat	199				
Arg Gln Asn Cys Gly Phe Pro Gly Val Thr Pro Ser Gln Cys Ala Asn					
	40 45 50				
aag ggc tgc tgt ttc gac gac acc gtt cgt ggg gtc ccc tgg tgc ttc	247				
Lys Gly Cys Cys Phe Asp Asp Thr Val Arg Gly Val Pro Trp Cys Phe					
	55 60 65				
tat cct aat acc atc gac gtc cct cca gaa gag gag tgt gaa ttt	292				
Tyr Pro Asn Thr Ile Asp Val Pro Pro Glu Glu Glu Cys Glu Phe					
	70 75 80				
tagacacttc tgcagggatc tgcctgcac ctgacggggt gccgtcccca gcacgggtgat	352				
tagtcccaga gtcgggctgc cacctccacc ggacacctca gacacgcttc tgcagctgtg	412				
cctcgggtca caacacagat tgactgctct gactttgact actcaaaatt ggcctaaaaa	472				
ttaaaagaga tcgatattaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa	532				
aaaaaaaaa	540				

FIG. 10

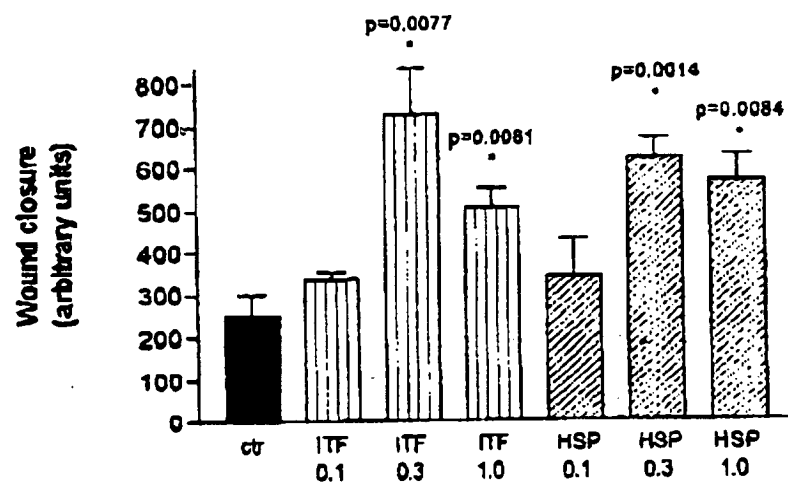


FIG. 11

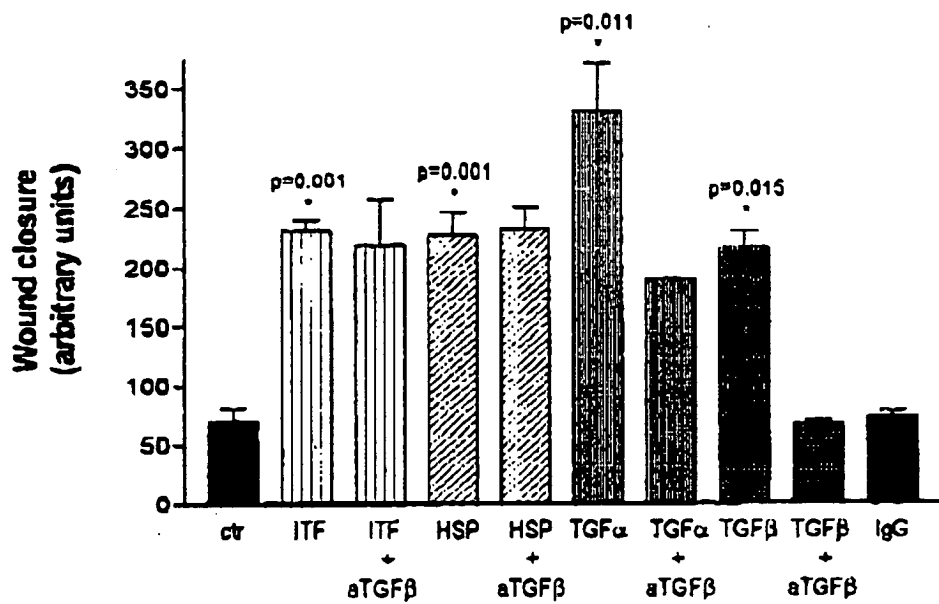


FIG. 12

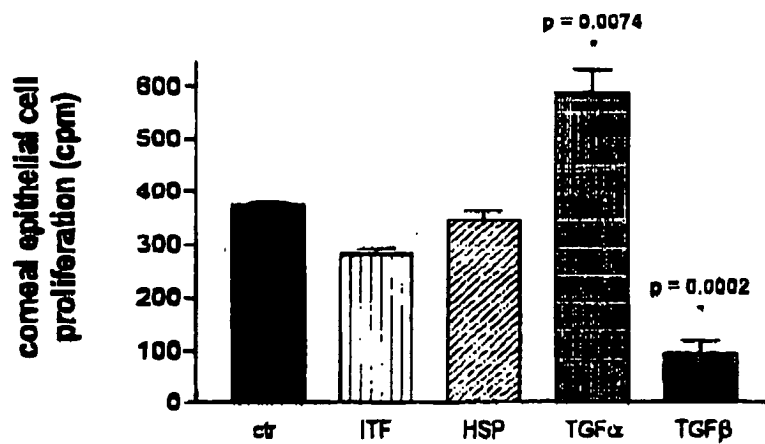


FIG. 13